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1: XM_171629. Homo sapiens simi...[gi:22062231]

Links

LOCUS LOC257238 1295 bp mRNA linear PRI 01-AUG-2002
DEFINITION Homo sapiens similar to cortical granule serine protease 1 precursor (LOC257238), mRNA.
ACCESSION XM_171629
VERSION XM_171629.1 GI:22062231
KEYWORDS
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 1295)
AUTHORS NCBI Annotation Project.
TITLE Direct Submission
JOURNAL Submitted (31-JUL-2002) National Center for Biotechnology Information, NIH, Bethesda, MD 20894, USA
COMMENT GENOME ANNOTATION REFSEQ: This model reference sequence was predicted from NCBI contig NT_009782 by automated computational analysis using gene prediction method: GenomeScan, supported by EST evidence.
 Also see:
 Documentation of NCBI's Annotation Process

FEATURES Location/Qualifiers
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 "organism="Homo sapiens"
 "db_xref="taxon:9606"
 "chromosome="17"
 protein 1..1295
 "gene="LOC257238"
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 "codon_start=1"
 "product="similar to cortical granule serine protease 1 precursor"
 "protein_id="XP_171629.1"

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1261 ctttatgttt tgtcatctta ctagcaacaa cataa

77

Revised: July 5, 2002.

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Document ID: 384

>XM_171629 ACCESSION:XM_171629 NID: gi 22062231 ref XM_171629.1 Homo sapiens similar to cortical granule serine protease 1 precursor (LOC257238), mRNA
Length = 1295

Identities = 141/153 (92%), Positives = 141 153 (92%)
Frame = +3

Query: 62 CGTAPLKDVLQGSFIIIGGTEAQAGAWPWVVSLSQIKYGRVLVHVEGGTLVRERWVLTAHC 121
CGTAPLKDVLQGSFIIIGGTEAQAGAWPWVVSLSQIKYGRVLVHVEGGTLVR
Sbjct: 3 CGTAPLKDVLQGSFIIIGGTEAQAGAWPWVVSLSQIKYGEVLVHVEGGTLVRE----- 155

Query: 122 TKDSDPLMWTAVIGTMNIHGFPYPHTKKIKIFATIHPNFILESYVNDIALFHLKKAVRYN 181
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Identities = 131/131 (100%), Positives = 131 131 (100%)
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Sequence # 0000004
File # 0000004
200 Human Proteins and their Substrates during the same

>XM_171629 ACCESSION:XM_171629 NID: gi 22062231 ref XM_171629.1 Homo sapiens similar to cortical granule serine protease 1 precursor (LOC257238), mRNA Length = 1295

Identities = 141/153 (92%), Positives = 141 153 (92%)
Frame = +3

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Identities = 131/131 (100%), Positives = 131/131 (100%)
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ALCFVILLATT

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